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OM protein protein search, using sw model

March 1, 2001, 15:47:01; Search time 210.42 Seconds (without alignments) 10.238 Million cell update

cell updates/sec

Title: Perfect score: US-09-331-631A-1_COPY_186_248 355

Sequence: KRDPQQREYEDCRRRCEQQE.....MMNPQRGGSGRYEEGEEEQS 63

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

268485 seqs, 34193795 residues

Total number of hits satisfying chosen parameters: 268485

Minimum DB

seq length: 0 length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database A_Geneseq_36:*

/SIDSi/gcgdata/geneseq/geneseqp/AA1980.DAT:*
/SIDSi/gcgdata/geneseq/geneseqp/AA1981.DAT:*
/SIDSI/gcgdata/geneseq/geneseqp/AA1982.DAT:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	10	20	21	18	10	19	13	19	19	19	19	19	:	
	P90996	Y55933	Y78914	W14783	P93109	W62837	R20181	W62831	W62832	W62830	W62829	W62828	ID	
•	Human androgen rec	Human ZC3 protein.	Human androgen rec	Androgen receptor.	Human androgen rec	Hordeum vulgare an	Sequence encoded b	-				Macadamia integrif	Description	

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13
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Mouse Wilms' tumou Human Wilms' tumou Human wilms' tumou Human metrastasis-a Protein regulating Transcription fact Mycobacterium spec Mycobacterium spec Mycobacterium snRNP 70K UI snRNP bindi 70K UI snRNP amino acid sequenc

Stenocarpus sinuat Chimeric Ewing's s Wilms' tumour poly Human WT1 protein Mouse WT1 protein

Mouse liver cancer Amino acid sequenc Zea mays antimicro HHV8 ORF 73 protei A human trichohyal Human androgen rec Human androgen rec

Peptide seq ID No: Epitope tagged TBP Protein of DNA mar SAP-AlaMet-VEGF165 SAP-GLYSer-VEGF165

G. max truncated Chimeric Ewing' T. gondii immunoge Human ovarian tumo

ALIGNMENTS

W62828 standard; Protein; 666 AA.

27-OCT-1998 (first entry)

Macadamia integrifolia antimicrobial protein

antimicrobial protein; infestation; control.

Macadamia integrifolia.

Key Protein Peptide /note= "signal peptide" 29..666 Location/Qualifiers

/note=

"mature protein"

WO9827805-A1

02-JUL-1998

22-DEC-1997; 97WO-AU00874

20-DEC-1996; 96AU-0004275.

(RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY

Bower NI, Goulter KC, Green JL, Manners JM, Marcus JP;

N-PSDB; V42310. WPI; 1998-377279/32.

0

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RESULT
W62829
ID W6
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Вb
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or mammals \ensuremath{\mathsf{N}}
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 34-36; 96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                      W62829;
                                                                                                                                                                                 20-DEC-1996;
                                                                                                                                                                                                                                            W09827805-A1
                                                                                                                                                                                                                                                                                                      Key
                                                                                                                                                                                                                                                                                                                        Macadamia
                                                                                                                                                                                                                                                                                                                                            antimicrobial protein; infestation; control
                                                                                                                                                                                                                                                                                                                                                              Macadamia integrifolia antimicrobial protein
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Sequence
                    animals.
                             The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and man
                                                           Claim 1; Page 39-41;
                                                                             Novel anti-microbial protein from e.g. Macadamia useful for controlling microbial infestations of
                                                                                                                                          Bower NI, Goulter KC,
                                                                                                                                                             (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
                                                                                                                                                                                                      22-DEC-1997;
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DB; V42311.
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 666
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 A
                                                                                                                                                                                  96AU-0004275
                                                                                                                                                                                                      97WO-AU00874.
                                                                                                                                                                                                                                                                        /note= "signal peptide"
29..666
                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                /note= "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                          96pp; English
                                                                                                                                          Green JL,
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Pred. No. 1.8e-32;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                          A
                                                                                                                                            Manners JM,
                                                                                                                                            Marcus
                                                                               integrifolia -
plants or mammals
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Query Match

93.5%;

Score 332;

DB 19;

Length

666;

RESULT

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Вb
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                                                                                          Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
Peptide
                                                                                                                                                      The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and majanimals.
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N-PSDB; V42316.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Macadamia integrifolia antimicrobial protein
                                                                                                                                   Sequence
                                                                                                                                                                                               Claim 1; Page 43-45;
                                                                                                                                                                                                                  Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or man
                                                                                                                                                                                                                                                                               Bower NI,
                                                                                                                                                                                                                                                                                                  (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
                                                                                                                                                                                                                                                                                                                      20-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      246 kgs 248
205 kgs
                                       145 krdpggreyedcrrhceggeprlqygcgrrcgeggrghgrggdlmnpgrggsgryeegee
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                   61 EQS 63
                                                  1 KRDPOQREYEDCRRRCEQQEPRQQHQCQLRCREQQQRQHGRGGDMMNPQRGGSGRYEEGEE 60
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1...28
                                                                                                                                                                                                                                                                                                                                                                                                      /note=
                                                                                          90.1%;
                                                                                                                                                                                               96pp; English
                                                                                                                                                                                                                                                                                                                                                                                                      "mature protein"
                                                                                                                                                                                                                                                                                Green
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4; Mismatches
                                                                                 Score 320; DB 19;
Pred. No. 1.5e-28;
4; Mismatches 3;
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                                                                                                                                                                                                                                                                                 Marcus
                                                                                                     Length 625;
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W62831
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Best Local :
                                                                                                                                   27-OCT-1998
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20-DEC-1996;
                  22-DEC-1997;
                                     02-JUL-1998
                                                      WO9827805-A1.
                                                                         Theobroma cacao
                                                                                            antimicrobial protein; infestation; control
                                                                                                              Theobroma cacao antimicrobial protein.
                                                                                                                                                                       W62831 standard; Protein; 525
                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-377279/32.
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                                                                                                                                                                                                                      141 kqqcvrecrekyqenpwrg--ereeeaeeeet 170
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                                                                                                                                                                                                                                          46
                                                                                                                                                                                                                                                          82 dpqrr-yeecqqecrqqeerqqpqcqqrclkrfeqeqqqsqrqfqecqqhchqqeqrpek 140
                                                                                                                                                                                                                                                                             3 DPQQREYEDCRRRCEQQEPRQQHQCQLRC----REQQ-----RQHGRGGDMM---
                                                                                                                                                                                                                                                                                                                                                                                                         Page 49-51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Goulter KC,
                                                                                                                                                                                                                                                                                                                                                 590 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                 -----NPQRGGSGRYEEGEEEQS 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein;
                  97WO-AU00874.
96AU-0004275
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                                                                                                                                                                                                                                                                                                         34.98;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infestation;
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                                                                                                                                                                                                                                                                                                  9;
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Pred. No. 1
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L.9e-06;
nes 15;
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plants or mammals
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RESULT
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Best Local :
The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocoa beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed clos
                                                                                    Claim 4; Fig 2; 59pp; English.
                                                                                                                      Recombinant cocoa proteins - are responsible for flavour in cocoa beans and produced in large quantities using yeast and bacterial
                                                                                                                                                              WPI; 1992-024418/03.
N-PSDB; Q20377.
                                                                                                                                                                                                Spencer ME,
                                                                                                                                                                                                                                                 11-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                             Theobroma cacao
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                                                                                                                                                                                                                                                                                                                                                                                                                                               R20181;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals
                                                                                                                                                                                                                         (MRSC ) MARS UK LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 KRDPQQREYEDCRRRCEQQ--EPRQQHQCQLRC----REQQRQHGRGGDMMNPQRGGSGR 54
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Pred. No. 4.9e-06;
3; Mismatches 17;
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Best Local Similarity 34.8%;
Matches 24; Conservative
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40.8%; Pred. No. 5.3e-06;
ative 13; Mismatches 17;
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Pred. No. 4.8e-05;
8; Mismatches 21;
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XX
AC P93109;
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DT 19-MAR-1990 (first entry)
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DE Human androgen receptor.
XX
KW Human androgen receptor; ployclo
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KW Human androgen receptor; ployclo
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KW Human androgen.
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PN W08909791-A.
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PD 19-OCT-1989; 89WO-US01548.
PF 13-APR-1989; 89WO-US01548.
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Best Local
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20-SEP-1996;
                                                                                          27-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Androgen receptor
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N-PSDB; N91772.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    French FS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-APR-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYNC-) UNIVERSITY OF NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 32.8 nes 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  58 qqqqqq--qqqqqqqqqqqqqqqqqqqqqqqqqqqqahrrgptgylvldeeqq 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 QQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wilson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
96WO-US15081.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    뙻
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 79; DB 1
Pred. No. 0.37;
S; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 10; Length 919;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22;
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RESULT
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Best Local
                                                                                                                                                                                                                                                                                                             Androgen receptor; AR; androgen-independent activation; inhibitor; cancer; benign prostatic hyperplasia; hirusutism; androgenic alopecia; acne; breast cancer; Kennedy disease; prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human androgen receptor (W14783) binds testosterone and, acting at the transcriptional level, regulates the growth of normal prostatic cells. Antisense oligonucleotides (see also T63200, T63404-05) based on an androgen receptor cDNA clone (see also T63407) can be used to prevent androgen receptor gene expression, thereby inhibiting the growth or survival of prostatic cells for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide(s) antisense to human androgen receptor and acidic FGF genes - used to inhibit gene expression, for the treatment of benign prostatic hyperplasia {\bf r}
           Disclosure; Page 7; 32pp; English
                                   screening compounds and such as prostate cancer
                                                            Novel non-androgen ligand binding peptides for inhibiting androgen-independent activation of androgen receptor, use
                                                                                                                                                       (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                 30-JUN-1998;
                                                                                                                                                                                                           30-JUN-1999;
                                                                                                                                                                                                                                      13-JAN-2000
                                                                                                                                                                                                                                                              WO200001813-A2
                                                                                                                                                                                                                                                                                                                                                                                             23-MAY-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y78914 standard; protein; 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WORC-) WORCESTER FOUND BIOMEDICAL RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 32.8 nes 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58 qqqqqq--qqqqqqqqqqqqqqetsprqqqqqqqedgspqahrrgptgylvldeeqq 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 QQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treatment of benign prostatic hyperplasia and prostate cancer
                                                                                                    2000-182113/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1997-202879/18
                                                                                                                              MD,
                                                                                                                             Bruchovsky N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      919 AA;
                                                                                                                                                                                 98US-0091871.
                                                                                                                                                                                                           99WO-CA00604.
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                                              and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22.3%;
                                              for treatment of androgen-mediated
                                                                                                                                                                                                                                                                                                                                                                    (AR) amino acid sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51pp; English.
                                                                                                                             Gout
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 79; DB 1
Pred. No. 0.37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA
                                                                                                                              PW,
                                                              androgen receptor, used
                                                                                                                              Snoek
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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                                                diseases
                                                               for
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC This sequence represents the human androgen receptor (AR) amino acid consequence. The invention relates to a fragment of the AR corresponding to amino acids 234-391 (see Y78913). The fragment mediates can acids 234-391 (see Y78913). The fragment mediates can acids 234-391 (see Y78913). The fragment mediates can accept the activation of the AR. The androgen receptor acts as can are activation of the AR. The androgen receptor acts as a transcription factor, regulating the expression of certain and activation pathway involves interaction with the androgen can activation pathway involves interaction with the androgen activation of the AR signal transduction pathway involves interaction with the androgen activation of the AR fragment and peptides derived from it can be used as agents for inhibiting androgen independent activation of the AR. The candidate activation domains, and as a tool for screening for compounds which affect androgen-independent activation of the AR. The peptides, when used in combination with androgen deprivation, effectively limit androgen mediated disease progression. These diseases include cancer, benign prostatic hyperplasia, hirusutism, androgenic alopecia, cane, breast cancer, Kennedy disease, and especially prostate cancer. The peptides and nucleic acids encoding them, are especially used for the treatment of androgen-mediated diseases, especially prostate tumours in cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                               myocadidata infarction, continuous and disorder; Parkinson's disease; amylotrophic lateral sclerosis; Leigh syndrome; cancer; cardiomyopathy; ischemic disorder; inflammation; diabetes mellitus; fibrosis; mitosis; mesandial disorder; growth regulation; wound healing; T cell activation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antirheumatic; antiathritic; antiinflammatory; antiallergic; osteopathic; antipsoriatic; antiatheriosclerotic; antiasthmatic; immunosupressive; neuroprotective; cardiant; cerebroprotect; cytostatic; antidiabetic; vulnery; STE20; protein kinase; STLK2; STLK3; STLK4; STLK5, STLK6, STLK7; ZC1, ZC2, ZC3, ZC4, KHS2, SULU1, SULU3, GEK2, PAK4; PAK5; antagonist; antibody; gene therapy; rheumatoid arthritis; artherosclerosis; asthma; inflammatory bowel disease; Crohn's disease; osteoarthritis; psoriasis;
                                         WPI; 1999-611301/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rhinitis; autoimmunity; organ transplantation; multiple sclerosis; myocardial infarction; cardiovascular disease; stroke; renal failure;
                                                                                  Plowman G,
                                                                                                                                                                         14-APR-1998;
                                                                                                                                                                                                                                                              21-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ZC3 protein
                                                                                                                                                                                                                  13-APR-1999;
                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 QQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ
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19; Conserv
                                                                                                                              SUGEN INC
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                                                                                    Martinez
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                         98US-0081784
                                                                                                                                                                                                                     99WO-US08150
                                                                                  ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22.3%; Score 79; DB 32.8%; Pred. No. 0.37 tive 15; Mismatches
                                                                                  Whyte
                                                                                    Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 21; Length 919, 0.37;
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N-PSDB; Z40485

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RESULT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Crohn's disease), asthma, osteoarthritis, psoriasis, atherosclerosis, rhinitis, autoimmunity, and organ transplantation, chronic inflammatory pelvic disease, multiple sclerosis, organ transplantation, myocardial infarction, cardiovascular disease, stroke, renal failure, oxidative stress-related neurodegenerative disorders (e.g. amylotrophic lateral sclerosis, Parkinson's disease and Leigh syndrome), cancer, cardiomyopathies, ischemic disorders, inflammatory disorders, diabetes mellitus, fibrotic and mesangial disorders. The proteins may also be useful for cell growth regulation (e.g. in wound healing), T cell activation, mitosis control, and as immunosuppressants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZC4, KHS2, SULU1, SULU3, GEK2, PAK4 and PAK5. The proteins are used to identify agonists and antagonists, and to raise antibodies. The polynucleotides are useful in gene therapy protocols. The polynucleotides, polypeptides, antibodies, antagonists and agonists may be used to treat diseases such as immune-related disorders and diseases (e.g. rheumatoid arthitis, artherosclerosis, chronic inflammatory bowel disease (e.g.
              (ARCH-) ARCH DEVELOPMENT CORP
                                                                                                                                                                                                                                                                                                                                                 Human androgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel kinase-related polypeptides used for the diagnosis and treatment of kinase-related diseases and disorders \,\cdot\,
                                           30-MAR-1988;
                                                                                                       05-OCT-1989
                                                                                                                                     W08909223-A
                                                                                                                                                                                                                                                                                                                     Androgen receptor;
                                                                        24-MAR-1989;
                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-1990
                                                                                                                                                                                                                                                                                                                                                                                                              P90996;
                                                                                                                                                                                                                                                                                                                                                                                                                                           P90996 standard; protein; 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       selected from STLK2,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   403 rreeerrqaerege 416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  QRGGSGRYEEGEEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           qrdpeahikhllhqrqrrieeqkeerrrveeqqrrereqrklqekeqqr---rledmqal 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 278-283; 387pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1326 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                           88US-0176107
                                                                                                                                                                                                                                                                                                                                                 receptor DNA clone
                                                                         89WO-US01238
                                                                                                                                                                                            /product=98 kD polypeptide 185..919
                                                                                                                                                                 /product=79 kD
                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                 /*tag≃
                                                                                                                                                                                                                                             ..919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to nucleic acid molecule encoding a kinase (2, STLK3, STLK4, STLK5, STLK6, STLK7, 2C1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -QR---EYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNP 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.3%;
                                                                                                                                                                                                                                                                                                                   polypeptide;
                                                                                                                                                                                                                             D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     novel STE20-related protein kinase. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 79; Pred. No.
                                                                                                                                                                 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      a kinase polypeptide
LK7, ZC1, ZC2, ZC3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
             This invention describes novel pure proapoptotic dependence peptides which comprise a sequence of an active dependence domain selected from dependence polypeptides consisting of p75NTR, androgen receptor, DCC, huntingtin polypeptide, Machado-Joseph disease gene product, SCA1, SCA2 SCA6 and atrophin-1 polypeptide. The proapoptotic peptides are capable of inducing cell death and can be used to develop products to mediate of
                                                                                                                                                                                                                                                                                                                                                                                         Proapoptotic; dependence domain; p75NTR; androgen receptor; DCC; huntingtin polypeptide; Machado-Joseph disease; SCA1; SCA2; SCA6; atrophin-1; cell death; apoptosis; Huntington's disease; head trauma; Alzheimer's disease; Kennedy's disease; spinocerebellar ataxia; stroke; dentatorubropallidolysian atrophy; cell proliferation; cell survival;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The protein is used to raise antibodies for receptor assays and affinity purification.

The 98 kD product starts at the first Met codon; the 79 kD product
                                                                                                       Disclosure; Page 178-179; 199pp; English
                                                                                                                               Bredesen DE,
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                                                                                                                                                                                                                         (BURN-) BURNHAM INST
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                                                                                                                                                                                                                                                   12-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                               neoplastic; malignant; autoimmune; fibrotic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       unliganded androgen receptor protein.
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19; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21.8%;
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can be used to develop products to mediate or hods can be used for reducing the severity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12;
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P;Ad. No. 0.55
12; Mismatches
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                     indicate that hAR is more closely related to glucocorticoid, mineralo-corticoid and progesterone receptors than to v-erb-A or receptors for oestrogen, vitamin D and thyroid hormones.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Spinocerebellar ataxias, dentatorubropallidoluysian atrophy, Machado-Joseph disease, stroke or head trauma. They can also be used for reducing the severity of a pathological condition mediated by upregulated cell proliferation or cell survival e.g. neoplastic, malignant, autoimmune or fibrotic conditions. This sequence represents a human unliganded androgen receptor described in the method of the invention.
                                                                        This sequence was deduced from a cDNA clone isolated by screening commercially available human testis and prostate lambda gt11 cDNA libraries. The sequence is very similar to that of rat AR and in the DNA-binding domain it is identical to that of rAR DNA-binding
                                                                                                                                                                         Androgen receptor and TR2 DNA binding proteins - DNA se and antibodies for detection and quantification methods
Sequence
                                                                                                                                                  Claim
                                                                                                                                                                                                                    N-PSDB; Q12001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a proapoptotic dependence domain mediated pathological conditions e.g. Huntington's disease, Alzheimer's disease, Kennedy's disease,
                                                                domain. Homology comparisons with other known steroid receptors
                                                                                                                                                                                                                                                                                      (ARCH-) ARCH DEV CORP
                                                                                                                                                                                                                                                                                                                 17-NOV-1989;
                                                                                                                                                                                                                                                                                                                                            19-OCT-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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2; Mismatches
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Best Local Similarity
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                                                                                                                                          a proapprotic dependence domain mediated pathological conditions e.g. Huntington's disease, Alzheimer's disease, Kennedy's disease, Spinocerebellar ataxias, dentatorubropallidoluysian atrophy. Machado-Joseph disease, stroke or head trauma. They can also be used for reducing the severity of a pathological condition mediated by upregulated cell proliferation or cell survival e.g. neoplastic, malignant, autoimmune or fibrotic conditions. This sequence represents a human androgen receptor described in the method of the invention.
                                                                                                                                                                                                                                                                           dependence polypeptides consisting of p75NTR, androgen receptor, DCC, huntingtin polypeptide, Machado-Uoseph disease gene product, SCA1, SCA2, SCA6 and atrophin-1 polypeptide. The proapoptotic peptides are capable of inducing cell death and can be used to develop products to mediate or inhibit apoptosis. The methods can be used for reducing the severity of
                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Proapoptotic; dependence domain; p75NTR; androgen receptor; DCC; huntingtin polypeptide; Machado-Joseph disease; SCA1; SCA2; SCA6; atrophin-1; cell death; apoptosis; Huntington's disease; head trauma; Alzheimer's disease; Kennedy's disease; spinocerebellar ataxia; stroke; dentatorubropallidoluysian atrophy; cell proliferation; cell survival;
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                                                                                                                                                                                                                                                                                                                                                                which comprise
                                                                                                                                                                                                                                                                                                                                                                                 This invention
                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                New proapoptotic dependence peptides, used to develop products for treating, e.g. Alzheimer's disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-561617/47
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QQREYEDCRRRCEQQEPRQQHQCQLRCREQQRQHGRGGDMMNPQRGGSGRYEEGEEEQ
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                                                                                                                   918
                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                 Page 90-93;
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                                                                                                                   AA;
                                                                                                                                                                                                                                                                                                                                                             describes novel pure proapoptotic dependence peptides a sequence of an active dependence domain selected from
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